> This Raw Listing contains the General Information
> Section and up to first 5 pages.

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 1   <110> APPLICANT: MORI, Satoshi
 2         NAKANISHI, Hiromi
 3         TAKAHASHI, Michiko
 4   <120> TITLE OF INVENTION: Nicotianamine Aminotransferase and Gene Therefor
 5   <130> FILE REFERENCE: 2185-0226P
 6   <140> CURRENT APPLICATION NUMBER: US/09/026,400
 7   <141> CURRENT FILING DATE: 1998-02-19
 8   <150> EARLIER APPLICATION NUMBER: 09-037499 JAPAN
 9   <151> EARLIER FILING DATE: 1997-02-21
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34         Arg Tyr Lys Ile Ser Ala Ser Val Glu Glu Ser Gly Pro Arg Pro Val
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36         ctg ccg ctg gcc cac ggt gac ccg tcc gtg ttc ccg gcc ttc cgc acg    349
37         Leu Pro Leu Ala His Gly Asp Pro Ser Val Phe Pro Ala Phe Arg Thr
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39         gcc gtc gag gcc gaa gac gcc gtc gcc gcc gcg ctg cgc acc ggc cag    397
40         Ala Val Glu Ala Glu Asp Ala Val Ala Ala Ala Leu Arg Thr Gly Gln
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42         ttc aac tgc tac gcc gcc ggc gtc ggc ctc ccc gcc gca cga agc gcc    445
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48    gac gtc ttc ctc acc gcc ggc gga act cag gcg atc gaa gtc ata atc    541
49    Asp Val Phe Leu Thr Ala Gly Gly Thr Gln Ala Ile Glu Val Ile Ile
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51    ccg gtg ctg gcc cag act gcc ggc gcc aac ata ctg ctt ccc cgg cca    589
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54    ggc tat cca aat tac gag gcg cga gcg gca ttc aac aag ctg gag gtc    637
55    Gly Tyr Pro Asn Tyr Glu Ala Arg Ala Ala Phe Asn Lys Leu Glu Val
56              180                 185                 190
57    cgg cac ttc gac ctc atc ccc gac aag ggg tgg gag atc gac atc gac    685
58    Arg His Phe Asp Leu Ile Pro Asp Lys Gly Trp Glu Ile Asp Ile Asp
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70    Glu Val Tyr Gly Lys Leu Val Leu Gly Ser Ala Pro Phe Ile Pro Met
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73    Gly Val Phe Gly His Ile Ala Pro Val Leu Ser Ile Gly Ser Leu Ser
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76    Lys Ser Trp Ile Val Pro Gly Trp Arg Leu Gly Trp Val Ala Val Tyr
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80    305                 310                 315                 320
81    aat tac ctt aat gtc tca acg gac cca gca acc ttc gtt cag gaa gct    1069
82    Asn Tyr Leu Asn Val Ser Thr Asp Pro Ala Thr Phe Val Gln Glu Ala
83              325                 330                 335
84    ctt cct aaa att ctt gag aac aca aaa gca gat ttc ttt aag agg att    1117
85    Leu Pro Lys Ile Leu Glu Asn Thr Lys Ala Asp Phe Phe Lys Arg Ile
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87    att ggt cta cta aag gaa tca tca gag ata tgt tat agg gaa ata aag    1165
88    Ile Gly Leu Leu Lys Glu Ser Ser Glu Ile Cys Tyr Arg Glu Ile Lys
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90    gaa aac aaa tat att acg tgt cct cac aag cca gaa gga tcg atg ttt    1213
91    Glu Asn Lys Tyr Ile Thr Cys Pro His Lys Pro Glu Gly Ser Met Phe
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93    gta atg gtc aaa cta aac tta cat ctt ttg gag gag atc cat gac gac    1261
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100      Pro Gly Ser Val Leu Gly Met Glu Asn Trp Val Arg Ile Thr Phe Ala
101                      420                 425                 430
102      tgc gtt cca tct tct ctt caa gat gga ctc gaa agg gtc aaa tca ttc      1405
103      Cys Val Pro Ser Ser Leu Gln Asp Gly Leu Glu Arg Val Lys Ser Phe
104                      435                 440                 445
105      tgt caa agg aac aag aag aag aat tct ata aat ggt tgt tagttgtaca       1454
106      Cys Gln Arg Asn Lys Lys Lys Asn Ser Ile Asn Gly Cys
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151          Gly Val Phe Gly His Ile Ala Pro Val Leu Ser Ile Gly Ser Leu Ser
152                      275                 280                 285
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154                      290                 295                 300
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203	His	Ala	Glu	Ala	Ala	Asp	Ala	Asn	Gly	Glu	Ser	Asn	Glu	His	Ala	Glu
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206	Asp	Ser	Ala	Ala	Asn	Gly	Glu	Ser	Asn	Gly	His	Ala	Ala	Ala	Ala	Ala
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209	Glu	Glu	Glu	Glu	Ala	Val	Glu	Trp	Asn	Phe	Ala	Gly	Ala	Lys	Asp	Gly
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212	Val	Leu	Ala	Ala	Thr	Gly	Ala	Asn	Met	Ser	Ile	Arg	Ala	Ile	Arg	Tyr
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215	Lys	Ile	Ser	Ala	Ser	Val	Gln	Glu	Lys	Gly	Pro	Arg	Pro	Val	Leu	Pro
216				160				165				170				
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234			255				260				265					
235	cca	aac	tac	gag	gcg	cgc	gcc	gcg	ttc	aac	agg	ctg	gag	gtc	cgg	cat	927
236	Pro	Asn	Tyr	Glu	Ala	Arg	Ala	Ala	Phe	Asn	Arg	Leu	Glu	Val	Arg	His
237			270				275				280					
238	ttc	gac	ctc	atc	ccc	gac	aag	ggg	tgg	gag	atc	gac	atc	gac	tcg	ctg	975
239	Phe	Asp	Leu	Ile	Pro	Asp	Lys	Gly	Trp	Glu	Ile	Asp	Ile	Asp	Ser	Leu
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241	gaa	tcc	atc	gcc	gac	aag	aac	acc	acc	gcc	atg	gtc	atc	ata	aac	ccc	1023
242	Glu	Ser	Ile	Ala	Asp	Lys	Asn	Thr	Thr	Ala	Met	Val	Ile	Ile	Asn	Pro
243				305				310				315				
244	aac	aac	ccg	tgc	ggc	agc	gtt	tac	tcc	tac	gac	cat	ctg	tcc	aag	gtc	1071

Please Note:
Use of n and/or Xaa have been detected in the Sequence Listing. Please review the
Sequence Listing to ensure that a corresponding explanation is presented in the <220> to
<223> fields of each sequence which presents at least one n or Xaa.

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